

Sequence alignment for 10/560,322

RESULT 1

US-09-850-948-2

; Sequence 2, Application US/09850948

; Patent No. 6919176

; GENERAL INFORMATION:

; APPLICANT: Yang, Jianxin

; APPLICANT: An, Songzhu

; APPLICANT: Tularik Inc.

; TITLE OF INVENTION: Polypeptides and Nucleic Acids Associated With Cancer

; FILE REFERENCE: 018781-008300US

; CURRENT APPLICATION NUMBER: US/09/850,948

; CURRENT FILING DATE: 2001-05-07

; NUMBER OF SEQ ID NOS: 29

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 2

; LENGTH: 362

; TYPE: PRT

; ORGANISM: Homo sapiens

; FEATURE:

; OTHER INFORMATION: human G-protein coupled receptor 4 (GPR4)

US-09-850-948-2

Query Match 98.7%; Score 1907; DB 2; Length 362;

Best Local Similarity 98.9%; Pred. No. 4.5e-185;

Matches 358; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

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Qy      1  MGNHTWEGCHVDSRVDHLHPPSLYIFVIGVGLPTNCLALWAAYRQVQQRNQLGVYLMNLS 60
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Db      1  MGNHTWEGCHVDSRVDHLFPSPSLYIFVIGVGLPTNCLALWAAYRQVQQRNELGVYLMNLS 60

Qy     61  IADLLYICTLPLWVDYFLHHDNWIHGPGSCKLPGFIFYTNIYISIAFLCCISVDRLAVA 120
      |||
Db     61  IADLLYICTLPLWVDYFLHHDNWIHGPGSCKLPGFIFYTNIYISIAFLCCISVDRLAVA 120

Qy    121  HPLRFARLRRVKTAVAVSSVWVWATELGANSAPLFHDELFRDRYNHTFCFEKFPMEGWVAV 180
      |||
Db    121  HPLRFARLRRVKTAVAVSSVWVWATELGANSAPLFHDELFRDRYNHTFCFEKFPMEGWVAV 180

Qy    181  MNLYRVFVGFLFPWALMLLSYRGILRAVRGSVSTERQEKAKIKRLALSLIAIVLVCFAPY 240
      |||
Db    181  MNLYRVFVGFLFPWALMLLSYRGILRAVRGSVSTERQEKAKIKRLALSLIAIVLVCFAPY 240

Qy    241  HVLLLSRSAIYLGRPWDGCFEERVFSAYHSSLAFTSLNLCVADPILYCLVNEGARSVDVAKA 300
      |||
Db    241  HVLLLSRSAIYLGRPWDGCFEERVFSAYHSSLAFTSLNLCVADPILYCLVNEGARSVDVAKA 300

Qy    301  LHNLLRFLASDKPQEMANASLTLETPLTSKRNSTAKAMTGSWAATPPSEGQVQLKMLPP 360
      |||
Db    301  LHNLLRFLASDKPQEMANASLTLETPLTSKRNSTAKAMTGSWAATPPSQGDQVQLKMLPP 360

Qy    361  AQ 362
      ||
Db    361  AQ 362
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Result 2

US-10-267-811-2

; Sequence 2, Application US/10267811
 ; Publication No. US20030109044A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Logan, Thomas Joseph
 ; APPLICANT: Galvin, Katherine M.
 ; TITLE OF INVENTION: METHODS OF USING 279, A HUMAN G
 ; TITLE OF INVENTION: PROTEIN-COUPLED PROTEIN RECEPTOR
 ; FILE REFERENCE: MPI01-227P1RM
 ; CURRENT APPLICATION NUMBER: US/10/267,811
 ; CURRENT FILING DATE: 2002-10-09
 ; PRIOR APPLICATION NUMBER: 60/329,648
 ; PRIOR FILING DATE: 2001-10-16
 ; NUMBER OF SEQ ID NOS: 3
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO 2
 ; LENGTH: 362
 ; TYPE: PRT
 ; ORGANISM: Homo sapien
 US-10-267-811-2

Query Match 98.7%; Score 1907; DB 4; Length 362;
 Best Local Similarity 98.9%; Pred. No. 5.1e-175;
 Matches 358; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy	1	MGNHTWEGCHVDSRV	DHLHPPSLYIFVIGVGLPTNCLALWAA	RQVQQRNQLGVYLMNLS	60
Db	1	MGNHTWEGCHVDSRV	DHLFPSPSLYIFVIGVGLPTNCLALWAA	RYRQVQQRNELGVYLMNLS	60
Qy	61	IADLLYICTLPLWVDYFLHHDNWIHGP	GSKLPGFIFYTNIYISIAFLCCISVDRYL	AVA	120
Db	61	IADLLYICTLPLWVDYFLHHDNWIHGP	GSKLPGFIFYTNIYISIAFLCCISVDRYL	AVA	120
Qy	121	HPLRFARLRRVKTAVAVSSV	VWATELGANSAPLFHDELFRDRYNHTFCFEK	FPMEGWVAV	180
Db	121	HPLRFARLRRVKTAVAVSSV	VWATELGANSAPLFHDELFRDRYNHTFCFEK	FPMEGWVAV	180
Qy	181	MNLYRVFVGFLFPWALMLLSYRGILRA	VRGSVSTERQEKAKIKRLALS	SLIAIVLVC	FAPY 240
Db	181	MNLYRVFVGFLFPWALMLLSYRGILRA	VRGSVSTERQEKAKIKRLALS	SLIAIVLVC	FAPY 240
Qy	241	HVLLLSRSAIYLGRP	WDCGFEERVFSAYHSSLAFTSLNCVADPILYCLVNEGARS	SDVAKA	300
Db	241	HVLLLSRSAIYLGRP	WDCGFEERVFSAYHSSLAFTSLNCVADPILYCLVNEGARS	SDVAKA	300
Qy	301	LHNLLRFLASDKPQEMANASLTLETPL	TSKRNSTAKAMTGSWAATPPSEGDQVQLKMLPP	360	
Db	301	LHNLLRFLASDKPQEMANASLTLETPL	TSKRNSTAKAMTGSWAATPPSQGDQVQLKMLPP	360	
Qy	361	AQ	362		
Db	361	AQ	36		

RESULT 3

A57641

G protein-coupled receptor 4 - human

C;Species: Homo sapiens (man)

C;Date: 08-Feb-1996 #sequence_revision 08-Feb-1996 #text_change 09-Jul-2004

C;Accession: A57641

R;Mahadevan, M.S.; Baird, S.; Bailly, J.E.; Shutler, G.G.; Sabourin, L.A.;

Tsilfidis, C.; Neville, C.E.; Narang, M.; Korneluk, R.G.

Genomics 30, 84-88, 1995

A;Title: Isolation of a novel G protein-coupled receptor (GPR4) localized to chromosome 19q13.3.
A;Reference number: A57641; MUID:96129306; PMID:8595909
A;Accession: A57641
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-362 <MAH>
A;Cross-references: UNIPROT:P46093; UNIPARC:UPI0000050428; GB:U21051;
NID:g687793; PIDN:AAA98457.1; PID:g687794
C;Genetics:
A;Gene: GDB:GPR4
A;Cross-references: GDB:371710; OMIM:600551
A;Map position: 19q13.3-19q13.3
A;Introns: #status absent
C;Superfamily: G protein-coupled receptor 4
C;Keywords: G protein-coupled receptor

Query Match 98.7%; Score 1907; DB 2; Length 362;
Best Local Similarity 98.9%; Pred. No. 1.6e-170;
Matches 358; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy	1	MGNHTWEGCHVDSRVDHLHPPSLYIFVIGVGLPTNCLALWAAYRQVQQRNQLGVYLMNLS	60
Db	1	MGNHTWEGCHVDSRVDHLFPSPSLYIFVIGVGLPTNCLALWAAYRQVQQRNELGVYLMNLS	60
Qy	61	IADLLYICTLPLWVDYFLHHDNWIHGPSCKLPGFIFYTNIYISIAFLCCISVDRYLAVA	120
Db	61	IADLLYICTLPLWVDYFLHHDNWIHGPSCKLPGFIFYTNIYISIAFLCCISVDRYLAVA	120
Qy	121	HPLRFARLRRVKTAVAVSSVWATELGANSAPLFHDELFRDRYNHTFCFEKFPMEGWAV	180
Db	121	HPLRFARLRRVKTAVAVSSVWATELGANSAPLFHDELFRDRYNHTFCFEKFPMEGWAV	180
Qy	181	MNLYRVFVGFLFPWALMLLSYRGILRAVRGSVSTERQEKAKIKRLALSIAIVLVCFAPY	240
Db	181	MNLYRVFVGFLFPWALMLLSYRGILRAVRGSVSTERQEKAKIKRLALSIAIVLVCFAPY	240
Qy	241	HVLLLSRSAIYLGRPWCDFEERVFSAYHSSLAFTSLNCVADPILYCLVNEGARSDVAKA	300
Db	241	HVLLLSRSAIYLGRPWCDFEERVFSAYHSSLAFTSLNCVADPILYCLVNEGARSDVAKA	300
Qy	301	LHNLLRFLASDKPQEMANASLTLETPLTSKRNSTAKAMTGSWAATPPSEGDQVQLKMLPP	360
Db	301	LHNLLRFLASDKPQEMANASLTLETPLTSKRNSTAKAMTGSWAATPPSQGDQVQLKMLPP	360
Qy	361	AQ	362
Db	361	AQ	362